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## **RAW SEQUENCE LISTING**

**The Biotechnology Systems Branch of the Scientific and Technical  
Information Center (STIC) no errors detected.**

Application Serial Number: 10/519,826  
Source: PCP/10  
Date Processed by STIC: 1/10/06

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PCT

**RAW SEQUENCE LISTING**  
**PATENT APPLICATION: US/10/519,826**

**DATE: 01/10/2006**  
**TIME: 11:49:55**

**Input Set : A:\BB-123.ST25.txt**  
**Output Set: N:\CRF4\01102006\J519826.raw**

3 <110> APPLICANT: Walther, Diego  
 4 Bader, Michael  
 6 <120> TITLE OF INVENTION: Neuronally Expressed Tryptophane Hydroxylase And its Use  
 8 <130> FILE REFERENCE: BB-123  
 10 <140> CURRENT APPLICATION NUMBER: 10/519,826  
 11 <141> CURRENT FILING DATE: 2004-12-30  
 13 <150> PRIOR APPLICATION NUMBER: PCT/EP03/07744  
 14 <151> PRIOR FILING DATE: 2003-07-16  
 16 <150> PRIOR APPLICATION NUMBER: DE 10232151.5  
 17 <151> PRIOR FILING DATE: 2002-07-16  
 19 <160> NUMBER OF SEQ ID NOS: 14  
 21 <170> SOFTWARE: PatentIn version 3.3  
 23 <210> SEQ ID NO: 1  
 24 <211> LENGTH: 2350  
 25 <212> TYPE: DNA  
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 33 aagaatatta caccgggatc catgcagcca gcaatgatga tgttttccag taaatactgg 180  
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 49 ttccctcgga agatctctga gttagacaaa tgctctcaca gagttctcat gtatggttct 660  
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 65 ccagacacat gccatgaact ctgggacat gttccactac ttgcggatcc taagttgct 1140  
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 75 accttccagg aagcctactt tgtttgcagaa agtttgcag aagccaaaga aaagatgagg 1440  
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181 Val Pro Gln Leu Glu Asp Val Ser Met Phe Leu Lys Glu Arg Ser Gly  
182 260 265 270  
185 Phe Thr Val Arg Pro Val Ala Gly Tyr Leu Ser Pro Arg Asp Phe Leu  
186 275 280 285  
189 Ala Gly Leu Ala Tyr Arg Val Phe His Cys Thr Gln Tyr Ile Arg His  
190 290 295 300  
193 Gly Ser Asp Pro Leu Tyr Thr Pro Glu Pro Asp Thr Cys His Glu Leu  
194 305 310 315 320  
197 Leu Gly His Val Pro Leu Leu Ala Asp Pro Lys Phe Ala Gln Phe Ser  
198 325 330 335  
201 Gln Glu Ile Gly Leu Ala Ser Leu Gly Ala Ser Asp Glu Asp Val Gln  
202 340 345 350  
205 Lys Leu Ala Thr Cys Tyr Phe Phe Thr Ile Glu Phe Gly Leu Cys Lys  
206 355 360 365  
209 Gln Glu Gly Gln Leu Arg Ala Tyr Gly Ala Gly Leu Leu Ser Ser Ile  
210 370 375 380  
213 Gly Glu Leu Lys His Ala Leu Ser Asp Lys Ala Cys Val Lys Ala Phe  
214 385 390 395 400  
217 Asp Pro Lys Thr Thr Cys Leu Gln Glu Cys Leu Ile Thr Thr Phe Gln  
218 405 410 415  
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222 420 425 430  
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255 aaataaggct atcaaaaagcg aggacaagaa aagccggcaaa gagcccgca aaggcgacac 300  
257 cacagagagg agcaagacag cagttgttt ctcccttgaag aatgaaggatg gtgggcttgt 360  
259 gaaaggactt agactattcc aggaaaaaca tgtcaacatg cttcatatcg aatccaggcg 420  
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265 tgagagcatt tggacggagg aagaagatct cgaggatgtg ccgtgggttcc ctcggaaagat 600  
267 ctctgagttt gacagatgtct ctcaccgagt cctcatgtac ggcaccggagc ttgtatgccc 660  
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281	cacccagtag	gtgcggcatg	gctccgaccc	cctctacacc	ccggaaccag	atacatgcca	1080									
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372 115 120 125  
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376 130 135 140  
379 Asp Leu Glu Asp Val Pro Trp Phe Pro Arg Lys Ile Ser Glu Leu Asp  
380 145 150 155 160  
383 Arg Cys Ser His Arg Val Leu Met Tyr Gly Thr Glu Leu Asp Ala Asp  
384 165 170 175  
387 His Pro Gly Phe Lys Asp Asn Val Tyr Arg Gln Arg Arg Lys Tyr Phe  
388 180 185 190  
391 Val Asp Val Ala Met Gly Tyr Lys Tyr Gly Gln Pro Ile Pro Arg Val  
392 195 200 205  
395 Glu Tyr Thr Glu Glu Glu Thr Lys Thr Trp Gly Val Val Phe Arg Glu  
396 210 215 220  
399 Leu Ser Lys Leu Tyr Pro Thr His Ala Cys Arg Glu Tyr Leu Lys Asn  
400 225 230 235 240  
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404 245 250 255  
407 Gln Leu Glu Asp Val Ser Met Phe Leu Lys Glu Arg Ser Gly Phe Thr  
408 260 265 270  
411 Val Arg Pro Val Ala Gly Tyr Leu Ser Pro Arg Asp Phe Leu Ala Gly  
412 275 280 285  
415 Leu Ala Tyr Arg Val Phe His Cys Thr Gln Tyr Val Arg His Gly Ser  
416 290 295 300  
419 Asp Pro Leu Tyr Thr Pro Glu Pro Asp Thr Cys His Glu Leu Leu Gly  
420 305 310 315 320  
423 His Val Pro Leu Leu Ala Asp Pro Lys Phe Ala Gln Phe Ser Gln Glu  
424 325 330 335  
427 Ile Gly Leu Ala Ser Leu Gly Ala Ser Asp Glu Asp Val Gln Lys Leu  
428 340 345 350  
431 Ala Thr Cys Tyr Phe Phe Thr Ile Glu Phe Gly Leu Cys Lys Gln Glu  
432 355 360 365  
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439 Leu Lys His Ala Leu Ser Asp Lys Ala Cys Val Lys Ser Phe Asp Pro  
440 385 390 395 400  
443 Lys Thr Thr Cys Leu Gln Glu Cys Leu Ile Thr Thr Phe Gln Asp Ala  
444 405 410 415  
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**VERIFICATION SUMMARY**

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